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us-09-488-265-27 x us-09-684-855-160 July 2, 2002 11:26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            title of invention: continuous fermentation process file reference: c38435/111692 . . .
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                  398 DRVVPLHGCGVDKLGRCKLDDFVEGLSFARSGGNWAECFA 437
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                                                                                                                                                                            298 DHTSTNHTLDSNPATFPLNATLYADESHDNTMVSIFFALGLYNGTKPLST 347
                                                                                                                                                                                                                                             251
                                                                                                                                                                                                                                                                                     248
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 51 LSRHGARYPTSSKSKKYSALIERIQKNATFKGKYAFLKTYNYTLGADDLT 100
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Length Weight:
                                                                                                                                                                                                                                                                                                                   NFTAVFAPPIRARLEALPGVNLTDEDVVNLMDMCPFDTVARTSDATQLSP 247
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DRVVPLHGCGVDKLGRCKLDDFVEGLSFARSGGNWAECFA 440
                                                                           TSVESIETDGYAASWTVPFAARAYVEMMQCEAGGGGGEGEKEPLVRVLVN 400
                                                                                                               TSVESIETDGYAASWTVPFAARAYVEMMQCEAGGGGGGGEGEKEPLVRVLVN 397
                                                                                                                                                         DHTSTNHTLDSNPATFPLNATLYADFSHDNTMVSIFFALGLYNGTKPLST 350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PFGENQMVNSGIKFYRRYKALARNIVPFVRASGSDRVIASAEKFIEGFQS 150
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5.304
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Average Mismatch: -2.248
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us-09-488-265-26 x us-09-273-871a-9 July 2, 2002 10:56
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Ratio: 4.949
Percent Similarity: 94.647
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Symbol comparison table: /SIDS/appl/gcg/gcgcore/data/rundata/blosum62.cmpCompCheck: 1102
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                               351 LNATLYADFSHDNTMVSIFFALGLYNGTKPLSTTSVESIEETDGYAASWT
                                                                                                                                                                                   351 LNATLYADFSHDNSMISIFFALGLYNGTAPLSTTSVESIEETDGYSASWT
                                                                                                                                                                                                                                                                                                                                      101 AIQKNATAFKGKYAFLKTYNYTLGADDLTPFGENQMVNSGIKFYRRYKAL 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 101 AIQKNATAFKGKYAFLKTYNYTLGADDLTPFGEQQMYNSGIKFYRRYKAL 150
451 EGLSFARSGGNWEECFA 467
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Length Weight:
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Ratio:
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Average Mismatch: -2.248
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GAP of: us-09-488-265-26 check: 9654 from: 1 to: 467

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us-09-488-265-26 x us-09-343-126b-167 July 2, 2002 10:57 ...
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Ratio: 5.315
Percent Similarity: 100.000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Symbol comparison table: /SIDS/appl/gcg/gcgcore/data/rundata/blosum62.cmp
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                                 351 LNATLYADFSHDNTMVSIFFALGLYNGTKPLSTTSVESIEETDGYAASWT 400
                                                                                                                                                                                                              301 KYYGYGAGNPLGPAQGYGFVNELIARLTHSPVQDHTSTNHTLDSNPATFP 350
                                                                                                                                                                                                                                                                                                                                                                                                                             125 ARKIVPFVRASGSDRVIASAEKFIEGFQSAKLADPGANPHQASPVINVII 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                    151 ARKIVPEVRASGSDRVIASAEKFIEGFQSAKLADPGANPHQASPVINVII 200
451 EGLSFARSGGNWEECFA 467
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Length Weight:
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                                                                                                                 LNATLYADFSHDNTMVSIFFALGLYNGTKPLSTTSVESIEETDGYAASWT 374
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Average Mismatch: -2.248
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GAP of: us-09-488-265-26 check: 9654 from: 1 to: 467

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Quality: 2448
Ratio: 5.242
Percent Similarity: 99.143
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                                                                                                                                                                                                                                    301 KYYGYGAGNPLGPAQGYGFVNELIARLTHSPVQDHTSTNHTLDSNPATFP
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                                                            401 VPFAARAYVEMMQCEAEKEPLVRVLVNDRVVPLHGCGVDKLGRCKRDDFV 450
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451 EGLSFARSGGNWEECFA 467
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Length Weight:
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                                                                                                                         LNATLYADFSHDNTMVSIFFALGLYNGTKPLSTTSVESIEETDGYSASWT 400
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                                        VPFAARAYVEMMQCEAEKEPLVRVLVNDRVVPLHGCGVDKLGRCKRDDFV 450
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Average Mismatch: -2.248
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GAP of: us-09-488-265-25

check: 291

from: 1

to: 1426

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Quality: 13940
Ratio: 9.929
Percent Similarity: 99.288
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence 25, application us/09488265
general information:
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451 acaaggctttggctagaaagattgttccattcgttagagcttctggttct 500
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us-09-488-265-27 x us-09-343-126b-167 July 2, 2002 11:24
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Ratio: 5.023
Percent Similarity: 97.907
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394 PLVRVLVNDRVVPLHGCGVDKLGRCKRDDFVEGLSFARSGGNWEECFA 441
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